

REMARKS

Applicant submits these Remarks and the sequence list, disk, and related information pursuant to the Examiner's Sequence Rule Letter (Paper 17).

Applicant, through undersigned counsel, has conferred with the Patent Office's staff at the Sequence Help Desk and believes that this submission corrects any errors identified. Applicant notes that many of the sequences disclosed are not of "genetic" origin, but rather are developed through the combinatorial metalloprotein library approach of the instant application.

Applicant notes that on the "Notice to Comply" a handwritten notation appears stating "SEQ ID'S for every sequence requiring compliance must be in spec." Applicant has submitted two amendments to the specification, one mailed on May 3, 2000 and one mailed on November 1, 2001. Applicant believes that between these two amendments all required disclosures have been made in the specification. If either amendment is not of record, or if Examiner has identified any error, the Examiner is requested to so inform Applicant.

Submitted herewith is the "Sequence Listing" identifiers amendment to the disclosure, wherein reference is made to the sequence by use of assigned identifier (37 CFR 1.821(d)); a substitute "Sequence Listing" computer readable form copy and substitute paper copy (37 CFR 1.821(e) and 1.824); and the required statement that the paper and computer readable copies are the same and include no new matter (37 CFR 1.821(g)).

In view of the above amendments and remarks, it is respectfully submitted that all grounds of objection to nucleotide sequence and/or amino acid sequence disclosures have been addressed. It is believed that the case is now in condition for examination and same is respectfully requested.

Authorization is given to charge payment of any additional fees required, or credit any overpayment, to Deposit Acct. 13-4213.

Respectfully submitted,

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